(b)   optionally, identifying amino acids in said (poly)peptide sequences to be modified so as to remove unfavorable interactions between amino acids within or between said or other (poly)peptide sequences;

(c)   identifying at least one structural sub-element within each of said (poly)peptide sequences;

(d)   backtranslating each of said (poly)peptide sequences into a corresponding coding nucleic acid sequence;

(e)   setting up cleavage sites in regions adjacent to or between the ends of sub-sequences encoding said sub-elements, each of said cleavage sites:

    (ea)   being unique within each of said coding nucleic acid sequences;

    (eb)   being common to the corresponding sub-sequences of any said coding nucleic acids;

wherein said vector is essentially devoid of any of said cleavage sites.

57.   A collection of vectors comprising a plurality of vectors according to claim 56.

58.   A vector according to claim 56, wherein said vector is an expression vector.

59.   A collection of vectors comprising a plurality of vectors according to claim 58.

60.   A kit comprising a vector according to claim 56.

61.   A kit comprising a collection of vectors according to claim 57.

62.   A kit comprising a vector according to claim 58.

63.   A kit comprising a collection of vectors according to claim 59.